

SEQUENCE LISTING PART OF THE DESCRIPTION

pONY8.1Z MLVHyb (SEQ ID NO 10)

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pONY8ZA CMVHyb (SEQ ID N 52)

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PEsynGP (SEQ ID No 53)

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PESDSYNGP (SEQ ID No 54)

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MLV construct CZCG (SEQ ID No 55)

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TACTTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGGTGATGCGGTTTTGGCA
GTACATCAATGGGCGTGGATAGCGGTTTGACTCACGGGGATTCCAAGTCTCCACCCCAT
30 TGACGTCAATGGGAGTTTGTTTTGGCACCAAAATCAACGGGACTTTCCAAATGTCTGTA
CAACTCCGCCCATTTGACGCAAAATGGGCGGTAGGCGTGTACGGTGGGAGGTCTATATAAG
CAGAGCTCGTTTAGTGAACGCGCCAGTCTTCCGATAGACTGCGTCGCCCCGGGTACCCGT
ATTCCCAATAAAGCCTCTTGCTGTTTGCATCCGAATCGTGGTCTCGCTGTTCTTGGGAG
GGTCTCCTCTGAGTGATTGACTACCCACGACGGGGGTCTTCATTTGGGGGCTCGTCCGG
35 GATTTGGAGACCCCTGCCAGGGACCACCGACCCACCACCGGGAGGTAAGCTGGCCAGCA
ACTTATCTGTGTCTGTCCGATTGTCTAGTGTCTATGTTTGTATGCGCCTGCGTCTG
TACTAGTTAGCTAACTAGCTCTGTATCTGGCGGACCCGTGGTGGAACTGACGAGTTCTGA
ACACCGGCGCGCAACCCTGGGAGAGCTCCAGGGACTTTGGGGGGCCGTTTTTGTGGCCCG
ACCTGAGGAAGGGAGTCGATGTGGAATCCGACCCCGTCAGGATATGTGGTTCTGGTAGGA
40 GACGAGAACCATAAACAGTTCCCGCCTCCGTCTGAATTTTGTCTTTCGGTTTGAACCGA
AGCCGCGCGTCTTGCTGTCTGCAGCGCTGCAGCATCGTTCTGTGTTGTCTGTCTGACT
GTGTTTCTGTATTTGTCTGAAAATTAGGGCCAGACTGTTACCACTCCCTTAAGTTTGACC
TTAGGTCACTGGAAGATGTGAGCGGATCGCTCACAACCAGTCGGTAGATGTCAAGAAG
AGACGTTGG

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PCGCLNGFR (SEQ ID No 57)

GTTACCTTCTGCTCTGCAGAATGGCCAACCTTTAACGTCGGATGGCCGCGAGACGGCACC
TTTAACCGAGACCTCATCACCCAGGTAAAGATCAAGGTCTTTTACCTGGCCCCGATGGA
50 CACCCAGACCAGGTCCCTACATCGTGACCTGGGAAGCCTTGGCTTTTGACCCCCCTCCC
TGGGTCAAGCCCTTTGTACACCCTAAGCCTCCGCTCCTCTTCTCCATCCGCCCCGTCT
CTCCCCCTTGAACCTCCTCGTTCGACCCCGCCTCGATCCTCCCTTTATCCAGCCCTCACT
CCTTCTCTAGGCGCCGGAATTCGTTAACTCGAGGATCCACCGGTGCGCCACCATGGTGAGC
AAGGGCGAGGAGCTGTTACCGGGGTGGTGCCCATCCTGGTTCGAGCTGGACGGCGACGTA
55 AACGGCCACAAGTTCAGCGTGTCCGGCGAGGGCGAGGGCGATGCCACCTACGGCAAGCTG
ACCCTGAAGTTCATCTGCACCACCGGCAAGCTGCCCGTGCCCTGGCCACCCTCGTGACC
ACCCTGACCTACGGCGTGCACTGCTTCAGCCGCTACCCCGACCACATGAAGACGACGAC
TTCTTCAAGTCCGCCATGCCCGAAGGCTACGTCCAGGAGCGACCATCTTCTTCAAGGAC
GACGGCAACTACAAGACCCGCGCGAGGTGAAGTTCGAGGGCGACACCCTGGTGAACCGC
60 ATCGAGCTGAAGGGCATCGACTTCAAGGAGGACGGCAACATCCTGGGGCACAAGCTGGAG

TACAACTACAACAGCCACAACGTCTATATCATGGCCGACAAGCAGAAGAACGGCATCAAG
GTGAACTTCAAGATCCGCCACAACATCGAGGACGGCAGCGTGCAGCTCGCCGACCACTAC
CAGCAGAACACCCCCATCGGCGACGGCCCCGTGCTGCTGCCCGACAACCACTACCTGAGC
ACCCAGTCCGCCCTGAGCAAAGACCCCAACGAGAAGCGCGATCACATGGTCTGTGGAG
5 TTCGTGACCGCCGCCGGGATCACTCTCGGCATGGACGAGCTGTACAAGTAAAGCGGCCCT
AGGGGTCTTTCCCCTCTCGCCAAAGGAATGCAAGGTCTGTTGAATGTCGTGAAGGAAGCA
GTTCTCTGGAAGCTTCTTGAAGACAAACAACGTCTGTAGCGACCCTTTGCAGGCAGCGG
AACCCCCACCTGGCGACAGGTGCCTCTGCGGCCAAAAGCCACCGAGTTGGTTCAGCTGC
TGCTGAGGCTGGACGACCTCGCGGAGTTCTACCGGCAGTGCAAATCCGTCGGCATCCAG
10 GAAACCAGCAGCGGCTATCCGCGCATCCATGCCCCGAAGTGCAGGAGTGGGGAGGCACG
ATGGCCGCTTTGGTCGAGGCGGATCCGGCCATTAGCCATATTATTCATTGGTTATATAGC
ATAAATCAATATTGGCTATTGGCCATTGCATACGTTGTATCCATATCATAATATGTACAT
TTATATTGGCTCATGTCCAACATTACCGCCATGTTGACATTGATTATTGACTAGTTATTA
ATAGTAATCAATTACGGGGTCATTAGTTCATAGCCCATATATGGAGTTCGCGCTTACATA
15 ACTTACGGTAAATGGCCCGCTGGCTGACCGCCCAACGACCCCGCCCATTGACGTCAAT
AATGACGTATGTTCCCATAGTAACGCCAATAGGGACTTTCCATTGACGTCAATGGGTGGA
GTATTTACGGTAAACTGCCCACTTGGCAGTACATCAAGTGTATCATATGCCAAGTACGCC
CCCTATTGACGTCAATGACGGTAAATGGCCCGCTGGCATTATGCCAGTACATGACCTT
ATGGGACTTTTCTACTTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGGTGAT
20 GCGGTTTTGGCAGTACATCAATGGGCGTGGATAGCGGTTTTGACTCACGGGATTTCCAAG
TCTCCACCCCAATTGACGTCAATGGGAGTTTGTTTTGGCACCAAAATCAACGGGACTTTCC
AAAATGTCTATAACAACCTCCGCCCATGACGCAATGGGCGGTAGGCATGTACGGTGGGA
GGTCTATATAAGCAGAGCTCGTTTGTGTAACCGTCAGATCGCCTGGAGACGCCATCCACG
CTGTTTTGACCTCCATAGAAGACACCGGGACCGATCCAGCCTCCGCGGCCCAAGCTTAC
25 CATGGGGGACGGTGCCACCGGCCGCGCATGGACGGGCGCGCCTGCTGCTGTTGCTGCT
TCTGGGGGTGTCCCTTGGAGGTGCCAAGGAGGCATGCCCCACAGGCCTGTACACACACAG
CGGTGAGTGCTGCAAAGCCTGCAACCTGGGCGAGGGTGTGGCCCAGCCTTGTGGAGCCAA
CCAGACCGTGTGTGAGCCCTGCCTGGACAGCGTGACGTTCTCCGACGTGGTGAGCGCGAC
CGAGCCGTGCAAGCCGTGCACCGAGTGCGTGGGGCTCCAGAGCATGTCCGCCCGTGCCT
30 GGAGGCCGACGACGCGGTGTGCCGTGCGCTACGGCTACTACCAGGATGAGACGACTGG
GCGCTCGAGGCGTGCCGCGTGTGCGAGCGGGCTCGGGCCTCGTGTTCTCTGCCAGGA
CAAGCAGAACACCGTGTGCGAGGAGTGCCCCGACGGCACGTATTCCGACGAGGCCAACCA
CGTGGACCCGTGCCTGCCCTGCACCGTGTGCGAGGACACCGAGCGCCAGCTCCGCGAGTG
CACACGCTGGGCGGACGCCGAGTGCGAGGAGATCCCTGGCCGTTGGATTACACGGTCCAC
35 ACCCCCAGAGGGCTCGGACAGCACAGCCCCAGCACCCAGGAGCCTGAGGCACCTCCAGA
ACAAGACCTCATAGCCAGCACGGTGGCAGGTGTGGTGACCACAGTGATGGGCAGCTCCCA
GCCCCGTGGTGACCCGAGGCACCACCGACAACCTCATCCCTGTCTATTGCTCCATCCTGGC
TGCTGTGGTGTGGGGCCTTGTGGCCTACATAGCCTTCAAGAGGTGGAACAGCTGCTGAGT
CGACTCTAGAGGATCCCCAACATCGATAAAATAAAAGATTTTATTTAGTCTCCAGAAAAA
40 GGGGGGAATGAAAGACCCACCTGTAGGTTTGGCAAGCTAGCTTAAGTAACGCCATTTTG
CAAGGCATGGAATAATACATAACTGAGAATAGAGAAGTTCAGATCAAGGTCAGGAACAGA
TGGAACAGCTGAATATGGGCCAAACAGGATATCTGTGGTAAGCAGTTCCTGCCCCGGCTC
AGGGCCAAGAACAGATGGAACAGCTGAATATGGGCCAAACAGGATATCTGTGGTAAGCAG
TTCTGCCCCGGCTCAGGGCCAAGAACAGATGGTCCCCAGATGCGGTCCAGCCCTCAGCA
45 GTTTCTAGAGAACCATCAGATGTTTCCAGGGTGCCCCAAGGACCTGAAATGACCCTGTGC
CTTATTTGAACTAACCAATCAGTTCGCTTCTCGCTTCTGTTTCGCGCGCTTCTGCTCCCCG
AGCTCAATAAAAGAGCCCAACCCCTCACTCGGGGCGCCAGTCCCTCCGATTGACTGAGT
CGCCCGGGTACCCGTGTATCCAATAAACCCCTTTCGAGTTGCATCCGACTTGTGGTCTCG
CTGTTCTTGGGAGGGTCTCCTCTGAGTGATTGACTACCCGTGAGCGGGGTCTTTTATT
50 TGGGGGCTCGTCCGGGATCGGGAGACCCCTGCCAGGGACCACCGACCCACCGGGGAG
GTAAGCTGGCTGCCTCGCGCGTTTTCGGTGATGACGGTGAAAACCTCTGACACATGCAGCT
CCCGGAGACGGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGCAGACAAGCCCGTCAGGG
CGCGTCAGCGGGTGTGGCGGGTGTGGGGGCGCAGCCATGACCCAGTCACGTAGCGATAG
CGGAGTGTATACTGGCTTAACTATGCGGCATCAGAGCAGATTGTACTGAGAGTGACCAT
55 ATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAAATACCGCATCAGGCGCTCTTCC
GCTTCTCGCTCACTGACTCGCTGCGCTCGGTGCTTTCGGCTGCGGCGAGCGGTATCAGCT
CACTCAAAGGCGGTAATACGGTTATCCACAGAATCAGGGGATAACGCAGGAAAGAACATG
TGAGCAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGCTTGTGGCGTTTTTTC
CATAGGCTCCGCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGA
60 AACCCGACAGGACTATAAAGATACCAGGCGTTTTCCCCTGGAAGCTCCCTCGTGCCTCT

CCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCCGCTTTCTCCCTTCGGGAAGCGTG
GCGCTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTTCGCTCCAAG
CTGGGCTGTGTGCACGAACCCCGTTACGCCCAGCGCTGCGCTTATCCGGTAACTAT
CGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAAC
5 AGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTCTTGAAGTGGTGGCCTAAC
TACGGCTACACTAGAAGGACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTC
GGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTT
TTTGGTTTGCAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATC
TTTTCTACGGGGTCTGACGCTCAGTGGAACGAAAACTCACGTTAAGGGATTTTGGTCATG
10 AGATTATCAAAAAGGATCTTCACCTAGATCCTTTTAAATTAATAATGAAGTTTAAATCA
ATCTAAAGTATATATGAGTAACTTGGTCTGACAGTTACCAATGCTTAATCAGTAGGCA
CCTATCTCAGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCTGACTCCCCGTCGTGTAG
ATAACTACGATACGGGAGGGCTTACCATCTGGCCCCAGTGCTGCAATGATACCGCGAGAC
CCACGCTCACCGGCTCCAGATTTATCAGCAATAAACAGCCAGCCGGAAGGGCCGAGCGC
15 AGAAGTGGTCTGCAACTTTATCCGCTCCATCCAGTCTATTAATTGTTGCCGGGAAGCT
AGAGTAAGTAGTTCCGCGAGTTAATAGTTTGCGCAACGTTGTTGCCATTGCTGCAGGCATC
GTGGTGTACGCTCGTCGTTTGGTATGGCTTCATTACGCTCCGGTTCCCAACGATCAAGG
CGAGTTACATGATCCCCATGTTGTGCAAAAAAGCGGTTAGCTCCTTCGGTCTCCGATC
GTTGTGCAAGTAAGTTGGCCGAGTGTTATCACTCATGGTTATGGCAGCACTGCATAAT
20 TCTCTTACTGTCTATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGAGTACTCAACCAAG
TCATTCTGAGAAATAGTGTATGCGGCGACCGAGTTGCTCTTGGCCGCGTCAATACGGGAT
AATACCGCGCCACATAGCAGAACTTTAAAGTGCTCATATTGGAAAACGTTCTTCGGGG
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CCCACTGATCTTCAGCATCTTTTACTTTTACCAGCGTTTCTGGGTGAGCAAAAACAGGA
25 AGGCAAAATGCCGCAAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTC
TTCTTTTTTCAATATTATTGAAGCATTATCAGGGTTATTGTCTCATGAGCGGATACATA
TTTGAATGTATTTAGAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCCGAAAAGTG
CCACCTGACGTCTAAGAAACCATTTATTCATGACATTAACCTATAAAAAATAGGCGTATC
ACGAGGCCCTTTTCGTCTCGCGCTTTTCGGTGATGACGGTGAAAACCTCTGACACATGCAG
30 TCCCCGGAGACGGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGCAGACAAGCCCGTCAG
GGCGGTCACGGGTGTTGGCGGGTGTGCGGGCTGGCTTAACTATGCGGCATCAGAGCAG
ATTGTACTGAGAGTGCACCATATGGACATATTGTCTGTAGAACGCGGCTACAATTAATAC
ATAACCTTATGTATCATAACATACGATTTAGGTGACACTATAGAAGTCTGACTCTAGAGT
CCGTTACATAACTTACGGTAAATGGCCCCGCTGGCTGACCGCCCAACGACCCCCGCCCAT
35 TGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTTTCCATTGACGTC
AATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATCAAGTGATCATATGC
CAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCCGCTGGCATTATGCCCAGT
ACATGACCTTATGGGACTTTTCTACTTTGGCAGTACATCTACGTATTAGTCATCGCTATTA
CCATGGTGATGCGGTTTTTGGCAGTACATCAATGGGCGTGGATAGCGGTTTGACTCACGGG
40 GATTTCCAAGTCTCCACCCCATTTGACGTCAATGGGAGTTTGTGTTGGCACCAAAATCAAC
GGGACTTTCCAAAATGTCGTAACAACTCCGCCCCATTGACGCAAATGGGCGGTAGGCGTG
TACGGTGGGAGGTCTATATAAGCAGAGCTCGTTTAGTGAACCGCGCCAGTCTTCCGATAG
ACTGCGTCGCCCCGGGTACCCGATTTCCCAATAAAGCCTCTTGCTGTTTGCATCCGAATCG
TGGTCTCGCTGTTTCTTGGGAGGGTCTCCTCTGAGTGATTGACTACCCACGACGGGGGTC
45 TTTCAATTGGGGGCTCGTCCGGGATTTGGAGACCCCTGCCAGGGACCACCGACCCACCA
CCGGGAGGTAAGCTGGCCAGCAACTTATCTGTGTCTGTCCGATTGTCTAGTGTCTATGTT
TGATGTTATGCGCTGCGTCTGTACTAGTTAGCTAACTAGCTCTGTATCTGGCGGACCCG
TGGTGGAAGTACGAGTTCTGAACACCCGCGCAACCTGGGAGACGTCCAGGGACTT
TGGGGGCGGTTTTTGTGGCCCGACCTGAGGAAGGGAGTCGATGTGGAATCCGACCCCGTC
50 AGGATATGTGGTTCTGGTAGGAGACGAGAACCTAAAACAGTTCCCGCCTCCGTCTGAATT
TTTGCTTTTCGGTTTGGAAACCGAAGCCGCGCTTGTCTGCTGCAGCGCTGCAGCATCGT
TCTGTGTTGTCTGTCTGACTGTGTTTCTGTATTTGTCTGAAAATTAGGGCCAGACTGT
TACCACTCCCTTAAGTTTGAACCTTAGGTCACTGGAAAGATGTCGAGCGGATCGCTCACAA
CCAGTCGGTAGATGTCAAGAAGAGACGTTGG

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PLTRI xP (SEQ ID No 58)

GCTAGCATAACTTCGTATAATGTATGCTATACGAAGTTATTCTAGAGAACCATCAGATGT
60 TTCCAGGGTGCCCCAAGGACCTGAAATGACCCGTGTCCTTATTTGAACTAACCAATCAGT

TCGCTTCTCGCTTCTGTTTCGCGCGCTTCTGCTCCCCGAGCTCAATAAAAGAGCCCACAAC
CCCTCACTCGGGCGCCAGTCCCTCCGATTGACTGAGTCGCCCAGGTACCCGTGTATCCAA
TAAACCTCTTGCAGTTGCATCCGACTTGTGGTCTCGCTGTTCTTGGGAGGGTCTCCTC
TGAGTGATTGACTACCCGTACGCGGGGGTCTTTTCAATTTGGGGGCTCGTCCGGGATCGGGA
5 GACCCCTGCCAGGGACCACCGACCCACCACCGGGAGGTAAGCTGGCTGCCTCGCGCGTT
TCGGTGATGACGGTGAAAACCTCTGACACATGCAGTCCCGGAGACGGTCACAGCTTGTG
TGTAAGCGGATGCCGGGAGCAGACAAGCCCGTCAGGGCGCGTCAGCGGGTGTGGCGGGT
GTCGGGGCGCAGCCATGACCCAGTCACGTAGCGATAGCGGAGTGTATACTGGCTTAAC
TGCGGCATCAGAGCAGATTGTACTGAGAGTGCACCATATGCGGTGTGAAATACCGCACAG
10 ATGCGTAAGGAGAAAATACCGCATCAGGCGCTCTTCCGCTTCCCTCGCTCACTGACTCGCT
GCGCTCGGTGCTTCGGCTGCGGCGAGCGGTATCAGCTCACTCAAAGGCGGTAATACGGTT
ATCCACAGAATCAGGGGATAACGCAGGAAAGAACATGTGAGCAAAAGGCCAGCAAAAGGC
CAGGAACCGTAAAAAGGCCGCTTGGCTGGCGTTTTTTCATAGGCTCCGCCCCCTGACGA
GCATCAGAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATA
15 CCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCCTCTCTGTTCCGACCCTGCCGCTTAC
CGGATACCTGTCCGCTTTCTCCCTTCGGGAAGCGTGGCGCTTCTCATAGCTCAGCGTG
TAGGTATCTCAGTTCCGGTGTAGGTGCTTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCC
CGTTCAGCCCGACCGCTGCGCCTTATCCGGTAACATATCGTCTTGAGTCCAACCCGGTAAG
ACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGT
20 AGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCTAACTACGGCTACACTAGAAGGACAGT
ATTTGGTATCTCGCTCTGCTGAAGCCAGTTACCTTCGGAAGAGTTGGTAGCTCTTG
ATCCGGCAAAACAAACACCGCTGGTAGCGGTGGTTTTTTTTGTTTGAAGCAGCAGATTAC
GCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCA
GTGGAACGAAAACTCACGTTAAGGGATTTTGGTCATGAGATTATCAAAAAGGATCTTAC
25 CTAGATCCTTTTAAATTAATAATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAAAC
TTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATT
TCGTTTCATCCATAGTTGCCTGACTCCCCGTGCTGTAGATAACTACGATACGGGAGGGCTT
ACCATCTGGCCCCAGTGCTGCAATGATACCGCGAGACCCACGCTACCGGCTCCAGATTT
ATCAGCAATAAACAGCCAGCCGGAAGGCGAGCGCAGAAGTGGTCTGCAACTTTATC
30 CGCTCCATCCAGTCTATTAATTGTTGCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAA
TAGTTTGCAGCAAGTTGTTGCCATTGCTGCAGGCATCGTGGTGTACGCTCGTCTGTTGG
TATGGCTTCATTACGCTCCGGTTCCCAACGATCAAGGCGAGTTACATGATCCCCATGTT
GTGCAAAAAGCGGTTAGCTCCTTCGGTCTCCGATCGTTGTCAGAAGTAAGTTGGCCGC
AGTGTATCACTCATGGTTATGGCAGCACTGCATAATTCTCTTACTGTCATGCCATCCGT
35 AAGATGCTTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTGAGAATAGTGTATGCG
GCGACCGAGTTGCTCTTGCCCGGCGTCAACACGGGATAATACCGCGCCACATAGCAGAAC
TTTAAAGTGCTCATCATTTGGAACGTTCTTCGGGGCGAAACTCTCAAGGATCTTACC
GCTGTTGAGATCCAGTTCGATGTAACCCACTCGTGCACCAACTGATCTTCAGCATCTTT
TACTTTACACAGCGTTTCTGGGTGAGCAAAAACAGGAAGGCAAAATGCCGCAAAAAGGG
40 AATAAGGGCGACACGGAAATGTTGAATACTCATACTCTTCTTTTCAATATTATTGAAG
CATTTATCAGGGTTATTGTCTCATGAGCGGATACATATTTGAATGTATTAGAAAAATAA
ACAAATAGGGGTTCGCGCACATTTCCCCGAAAAGTGCCACCTGACGTCTAAGAAACCAT
TATTATCATGACATTAACCTATAAAAATAGGCGTATCAGAGGCCCTTTCGTCTTCAAGA
ATTCATACCAGATCACCGAAAATGTCTCCAAATGTGTCCCCCTCACACTCCCAAATTC
45 GCGGGCTTCTGCCTCTTAGACCACTCTACCCTATTCCCCACACTACCGGAGCCAAAGCC
GCGGCCCTTCGTTTTCTTTGCTTTTGAAAGACCCACCCGTAGGTGGCAA

LTR plasmid (SEQ ID No 59)

GCTAGCTTAAGTAACGCCATTTTGCAAGGCATGGAAAAATACATAACTGAGAATAGAGAA
50 GTTCAGATCAAGGTCAGGAACAGATGGAACAGCTGAATATGGGCCAAACAGGATATCTGT
GGTAAGCAGTTCTGCCCCGGCTCAGGGCCAAGAACAGATGGAACAGCTGAATATGGGCC
AAACAGGATATCTGTGGTAAGCAGTTCTGCCCCGGCTCAGGGCCAAGAACAGATGGTCC
CCAGATGCGGTCCAGCCCTCAGCAGTTCTAGAGAACCATCAGATGTTTCCAGGGTGCC
CAAGGACCTGAAATGACCCTGTGCCTTATTTGAACTAACCAATCAGTTCGCTTCTCGCTT
55 CTGTTTCGCGCGCTTCTGCTCCCCGAGCTCAATAAAAGAGCCCACAACCCCTCACTCGGG
CGCCAGTCTCCGATTGACTGAGTCGCCCAGGTACCCGTGTATCCAATAAACCCCTCTTG
AGTTGCATCCGACTTGTGGTCTCGCTGTTCCTTGGGAGGGTCTCCTCTGAGTGATGACT
ACCCGTACGCGGGGGTCTTTCAATTTGGGGGCTCGTCCGGGATCGGGAGACCCCTGCCAG
GGACCACCGACCCACACCGGGAGGTAAGCTGGCTGCCTCGCGCTTTCGGTGATGACGG

TGAAAACCTCTGACACATGCAGCTCCCGGAGACGGTCACAGCTTGTCTGTAAGCGGATGC
CGGGAGCAGACAAGCCCGTCAGGGCGCGTCAGCGGGTGTGGCGGGTGTGGGGCGCAGC
CATGACCCAGTCACGTAGCGATAGCGGAGTGTATACTGGCTTAACATGCGGCATCAGAG
CAGATTGTACTGAGAGTGCACCATATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGA
5 AAATACCGCATCAGGCGCTCTCCGCTTCCTCGCTCACTGACTCGCTGCGCTCGGTCTGTT
CGGCTGCGGCGAGCGGTATCAGCTCACTCAAAGGCGGTAATACGGTTATCCACAGAATCA
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AAGGCCGCGTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAAAT
CGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGGCGTTTCCC
10 CCTGGAAGCTCCCTCGTGCGCTCTCCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCC
GCCTTTCTCCCTTCGGGAAGCGTGGCGCTTTCTCATAGCTCACGCTGTAGGTATCTCAGT
TCGGTGTAGGTCTGTTCCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCCGTTACGCCCGAC
CGCTGCGCCTTATCCGGTAACATATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCG
CCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACA
15 GAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGGACAGTATTTGGTATCTGC
GCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAACAA
ACCACCGCTGGTAGCGGTGGTTTTTTTTGTTTGAAGCAGCAGATTACGCGCAGAAAAAA
GGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGAACGAAAAAC
TCACGTTAAGGGATTTTGGTTCATGAGATTATCAAAAAGGATCTTCACCTAGATCCTTTTA
20 AATTAAAAATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAACTTGGTCTGACAGT
TACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTCTGTTTCATCCATA
GTTGCCTGACTCCCCGTCGTGTAGATAACTACGATACGGGAGGGCTTACCATCTGGCCCC
AGTGCTGCAATGATACCGCGAGACCCACGCTCACC GGCTCCAGATTTATCAGCAATAAAC
CAGCCAGCCGGAAGGGCCGAGCGCAGAAGTGGTCTGCAACTTTATCCGCCTCCATCCAG
25 TCTATTAATTGTTGCCGGAAGCTAGAGTAAGTAGTTCCCGAGTTAATAGTTTGGCGAAC
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30 GTGACTGGTGAGTACTCAACCAAGTCATTCTGAGAATAGTGTATGCGGCGACCGAGTTGC
TCTTGCCCCGGCTCAACACGGGATAATACCGCGCCACATAGCAGAACTTTAAAAGTGCTC
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35 CGGAAATGTTGAATACTCATACTCTTCTTTTTCAATATTATTGAAGCATTATCAGGGT
TATTGTCTCATGAGCGGATACATATTTGAATGTATTTAGAAAAATAAACAATAGGGGT
CCGCGCACATTTCCCCGAAAAGTGCCACCTGACGTCTAAGAAACCATTATTATCATGACA
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CACC GAAAACGTCTCCTCAAATGTGTCCCCCTCACACTCCCAAATTCGCGGGCTTCTGCC
40 TCTTAGACCACTCTACCCTATTCCCCACACTCACC GGAGCCAAAGCCGCGGCCCTTCCGT
TTCTTTGCTTTTGAAGACCCCAACCGTAGGTGGCAA